

SEQUENCE LISTING

10/510677

SEQ ID NO.: 1; AAC2-1 nucleotide sequence

5 AC
 TCGCCACTCCTCCGACGTGCTGGGCAACCTCAACGAGCTGCGCCTGCGCGGGATCCTCACTGACGTCACGCTGCT
 GG
 TTGGCGGGCAACCCCTCAGAGCACACAAGGCAGTTCTCATCGCCTGCAGTGGCTTCTTCTATTCAATTTTCCGGG
 GC
 10 CGTGCGGGAGTCGGGGTGGACGTGCTCTCTCTGCCCGGGGGTCCCGAAGCGAGAGGCTTCGCCCCCTCTATTGGAC
 TT
 CATGTACACTTCGCGCCTGCGCCTCTCTCCAGCCACTGCACCAGCAGTCCTAGCGGCCGCCACCTATTTGCAGAT
 GG
 AGCACGTGGTCCAGGCATGCCACCGCTTCATCCAGGCCAGCTATGAACCTCTGGGCATCTCCCTGCGCCCCCTGG
 15 AA
 GCAGAACCCCCAACACCCCCAACGGCCCCCTCCACCAGGTAGTCCAGGCGCTCCGAAGGACACCCAGACCCACCT
 AC
 TGAATCTCGAAGCTGCAGTCAAGGCCCCCAGTCCAGCCAGCCCTGACCCCAAGGCCTGCAACTGGAAAAAGTA
 CA
 20 AGTACATCGTGCTAAACTCTCAGGCCTCCCAAGCAGGGAGCCTGGTTCGGGGAGAGAAGTTCTGGTCAACCTTGCC
 CC
 CAAGCCAGGCTCCCCAGTGGAGACGAGGCCTCCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGTGAAGAAGGA
 CC
 CATTCCTGGTCCCCAGAGCAGGCTCTCTCCAAGTGTGCCACTGTGCAGTTCAAATGTGGGGCTCCAGCCAGTAC
 25 CC
 CCTACCTCCTCACATCCCAGGCTCAAGACACCTCTGGATCACCTCTGAACGGGCTCGTCCACTACCGGGAGTGA
 AT
 TTTTCAGCTGCCAGAAGTGTGAGGCTGTGGCAGGGTGCTCATCGGGGGCTGGACTCCTTGGTTCTGGGGACGAA
 GA
 30 CAAACCTATAAGTGTGAGCTGTGCCGGTCTTCGTTCCGCTACAAGGGCAACCTTGCCAGTCACCGTACAGTGCA
 CA
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 GC
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 35 CA
 CGTGCTGATCCACACCGGGGAGAAGCCCTACCCTTGCCCTACCTGCGGAACCCGCTTCCGCCACCTGCAGACCCCT
 CA
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 GT
 40 CAACTGCGGCTGCATCTGCGCCAGAAACACGGAGCTGTACCAACACCAAAGTGCACTACCACATTCTCGGGGGG
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 TC
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 45 GC
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 50 TGGCCCCCATTGCATTGAGTTTATCTGTAAAATATAATTTATTGAGGCCTTTGGGTGGCACCAGGGGCTTCATTC
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 GC
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 55 TT
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10 GT
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15 TAACCCATCCTTTACTACAGAGGCATATGGGTTTGAATGTTACCTGGGGTTCTCTCTATTGAGTTGAGCCCCCTC
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20 AT
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25 TAAGAGGTTGGTTGAGGGGTGCAGTTTCTGGTGTAGGCCAGGTAGGTAGAAAGTGAGGAACAGGGTTGCCTCTTG
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30 TT
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GC
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35 GCAGGTGGAGAATGCCTGGGGGTAGAAATGTTAGATCTTGCAACATCAGATCCTTGAATAAAGAAGCCTCTCTG
CG
CAAAAAAAAAAAAAAAAAAAAAA

SEQ ID NO.: 2; AAC2-2 Open reading frame

ATGGGTTCCCCCGCGCCCGGAGGGAGCGCTGGGCTACGTCCGAGTTCACTCGCCACTCCTCCGACGTGCTGGGCAACCT
CAACGAGCTGCGCCTGCGCGGGATCCTCACTGACGTACGCTGCTGGTTGGCGGGCAACCCCTCAGAGCACACAAGGCAGTTC
5 TCATCGCCTGCAGTGGCTTCTTCTATTCAATTTTCCGGGGCCGTGCGGGAGTCGGGGTGGACGTGCTCTCTCTGCCCCGGGGT
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CCTAGCGGCGCCACCTATTTGCAGATGGAGCACGTGGTCCAGGCATGCCACCGCTTCATCCAGGCCAGCTATGAACCTCTGG
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10 CACCCAGACCCACCTACTGAATCTCGAAGCTGCAGTCAAGGCCCCCAGTCCAGCCAGCCCTGACCCCAAGGCCTGCAACTG
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15 ACTGTGAGGCTGTGGCAGGGTGTCTATCGGGGCTGGACTCCTTGGTTCTGGGGACGAAGACAAACCCTATAAGTGTCACTG
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20 CCCCTGTGGCCTGCATTTCGGGCACAAGAGTCAACTGCGGCTGCATCTGCGCCAGAAACAGGAGCTGCTACCAACACCAAAG
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SEQ ID NO.: 3; 7524

ATACCCGGAACCTCCCTAAGCCTTCTATTAGCTCCAATAATAGTAAGCCTGTGGAAGACAAAGATG

25 SEQ ID NO.: 4; 7526

GCCTGTGTCCCCTAGACTCCAACCTCAGCAACGGAAATAGAACTCTGACCCTGTTTAACGTGACCAGGAAC

SEQ ID NO.: 5; 7528

30 ACGTGCTTTACGGACCCGATGCTCCTACAATCAGCCCTCTAAACACAAGCTATAGATCAGGGGAAAATCT

SEQ ID NO.: 6; 7533

ACGTAAACAGGGTCAGAGTTCTATTTCCGTTGCTGAGTTGGAGTCTAGGGGACACAGGCAGGGACTGGT

SEQ ID NO.: 7; 7535

35 CTGATCTATAGCTTGTGTTTAGAGGGCTGATTGTAGGAGCATCGGGTCCGTAAAGCACGTTGAGAATCAC

SEQ ID NO.: 8; 7537

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40 SEQ ID NO.: 9; 7567

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SEQ ID NO.: 10; 7568

45 TGAATCAGACCTCCTGGCGCTGACTGGATTTTGGGTTTCGCATTTGTAGCTTGCTGTGTCGTTCCCTGGTC

SEQ ID NO.: 11; 7576

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50 SEQ ID NO.: 12; 7587

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SEQ ID NO.: 13; 7677

55 TTCTAACCAGTTGAGGATGAGGACGCAGTTGCATTAACTTGTGAGCCAGAGATTCAAAATACCACTTATTTATG
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SEQ ID NO.: 14; 7678

60 GTCTAATGATAACCGCACATTGACACTCCTGTCCGTTACTCGCAATGATGTAGGACCTTATGAGTGTGGCATTCA
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SEQ ID NO.: 15; 7679

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TTGCC

5 SEQ ID NO.: 16; 7680

TGATGGAAACATTTCAGCAGCATACTCAAGAGTTATTTATAAGCAACATAACTGAGAAGAACAGCGGACTCTATAC
TTGCC

10 SEQ ID NO.: 17; 7681

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15 SEQ ID NO.: 18; 7682

ATGTGCGGTTATCATTAGACAACCTGCAAGCGTGGGCTAACCGGCAAACCTTTGGTTATTGACCCACCATAAATAAG
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20 SEQ ID NO.: 19; 7683

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ACTCA

25 SEQ ID NO.: 20; 7684

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AGTTC

30 SEQ ID NO.: 21; 7685

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35 SEQ ID NO.: 22; 7686

CCTCAGGTTTCACAGGTGAAGGCCACAGCATCCTTGTCTCCACGGGT

SEQ ID NO.: 23; CEA-CAP6D

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ACAGCCTCAC TTCTAACCTT CTGGAACCCG CCCACCACTG CCAAGCTCAC TATTGAATCC
35 ACGCCGTTCA ATGTCGCAGA GGGGAAGGAG GTGCTTCTAC TTGTCCACAA TCTGCCCCAG
CATCTTTTTG GCTACAGCTG GTACAAAGGT GAAAGAGTGG ATGGCAACCG TCAAATTATA
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ATATACCCCA ATGCATCCCT GCTGATCCAG AACATCATCC AGAATGACAC AGGATTCTAC
ACCCATACAG TCATAAAGTC AGATCTTGTC AATGAAGAAG CAACTGGCCA GTTCCGGGTA
40 TACCCGAGC TGCCCCAAGCC CTCCATCTCC AGCAACAAC CCAAACCCGT GGAGGACAAG
GATGCTGTGG CTTTACCTG TGAACCTGAG ACTCAGGACG CAACCTACCT GTGGTGGGTA
AACAATCAGA GCCTCCCGGT CAGTCCCAGG CTGCAGCTGT CCAATGGCAA CAGGACCCCTC
ACTCTATTCA ATGTCACAAG AAATGACACA GCAAGCTACA AATGTGAAAC CCAGAACCCA
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45 ACCATTTCCC CTCTAAACAC ATCTTACAGA TCAGGGGAAA ATCTGAACCT CTCCTGCCAC
GCAGCCTCTA ACCACCTGC ACAGTACTCT TGGTTTGTCA ATGGGACTTT CCAGCAATCC
ACCCAAGAGC TCTTTATCCC CAACATCACT GTGAATAATA GTGGATCCTA TACGTGCCAA
GCCATAACT CAGACACTGG CCTCAATAGG ACCACAGTCA CGACGATCAC AGTCTATGAG
CCACCCAAAC CTTTCATCAC CAGCAACAAC TCCAACCCCG TGGAGGATGA GGATGCTGTA
50 GCCTTAACCT GTGAACCTGA GATTCAGAAC ACAACCTACC TGTGGTGGGT AAATAATCAG
AGCCTCCCGG TCAGTCCCAG GCTGCAGCTG TCCAATGACA ACAGGACCCCT CACTCTACTC
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GACCACAGCG ACCCAGTCAT CCTGAATGTC CTCTATGGCC CAGACGACCC CACCATTTC
CCCTCATACA CCTATTACCG TCCAGGGGTG AACCTCAGCC TCTCCTGCCA TGCAGCCTCT
55 AACCACCTG CACAGTATTC TTGGCTGATT GATGGGAACA TCCAGCAACA CACACAAGAG
CTCTTTATCT CCAACATCAC TGAGAAGAAC AGCGGACTCT ATACCTGCCA GGCCAATAAC
TCAGCAGTG GCCACAGCAG GACTACAGTC AAGACAATCA CAGTCTCTGC GGAGCTGCCC
AAGCCCTCCA TCTCCAGCAA CAACTCCAAA CCCGTGGAGG ACAAGGATGC TGTGGCCTTC
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60 CCAGTCAGTC CCAGGCTGCA GCTGTCCAAT GGCAACAGGA CCCTCACTCT ATTCAATGTC

5 ACAAGAAATG ACGCAAGAGC CTATGTATGT GGAATCCAGA ACTCAGTGAG TGCAAACCGC
 AGTGACCCAG TCACCCTGGA TGTCTCTAT GGGCCGGACA CCCCATCAT TTCCCCCCA
 GACTCGTCTT ACCTTTCGGG AGCGGACCTC AACCTCTCCT GCCACTCGGC CTCTAACCCA
 TCCCCGAGT ATTCTTGGCG TATCAATGGG ATACCGCAGC AACACACACA AGTTCTCTTT
 ATCGCCAAAA TCACGCCAAA TAATAACGGG ACCTATGCCT GTTTTGTCTC TAACTTGGCT
 ACTGGCCGCA ATAATTCCAT AGTCAAGAGC ATCACAGTCT CTGCATCTGG AACTTCTCCT
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SEQ ID NO.: 24; CAP6D-1,2

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 15 ATATACCCCA ATGCATCCCT GCTGATCCAG AACATCATCC AGAATGACAC AGGATTCTAC
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 25 GCTCACAATA GCGACACCGG ACTCAACCGC ACAACCGTGA CGACGATTAC CGTGTATGAG
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 30 GATCACTCCG ACCCTGTTAT CCTTAATGTT TTGTATGGCC CAGACGACCC AACTATATCT
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 40 GACTCGTCTT ACCTTTCGGG AGCGGACCTC AACCTCTCCT GCCACTCGGC CTCTAACCCA
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 45